

FIGURE 1.

A

CTCTAAGGTTTCGGGGTGAATCCTTGGGCCGCTGGGCAAGCGGCGAGACCTGGCCAGGGCCAGCGAGCCGAGGACAGAGGGCGCACGGAGGGCCGGG 100  
 CGCAGCCCCGGCGCTTGCAGACCCCGCC ATG GAC CCG TTC CTG GTG CTG CTG CAC TCG GTG TCG TCC AGC CTG TCG AGC  
 Met Asp Pro Phe Leu Val Leu Leu His Ser Val Ser Ser Ser Leu Ser Ser

200 Clone 15 Clone 8  
 AGC GAG CTG ACC GAG CTC AAG TTC CTA TGC CTC GGG CGC GTG GGC AAG CGC AAG CTG GAG CGC GTG CAG AGC GGC  
 Ser Glu Leu Thr Glu Leu Lys Phe Leu Cys Leu Gly Arg Val Gly Lys Arg Lys Leu Glu Arg Val Gln Ser Gly 42

300  
 CTA GAC CTC TTC TCC ATG CTG CTG GAG CAG AAC GAC CTG GAG CCC GGG CAC ACC GAG CTC CTG CGC GAG CTG CTC  
 Leu Asp Leu Phe Ser Met Leu Leu Glu Gln Asn Asp Leu Glu Pro Gly His Thr Glu Leu Leu Arg Glu Leu Leu 67

400  
 GCC TCC CTG CGG CGC CAC GAC CTG CTG CGG CGC GTC GAC GAC TTC GAG GCG GGG GCG GCG GCC GGG GCC GCG CCT  
 Ala Ser Leu Arg Arg His Asp Leu Leu Arg Arg Val Asp Asp Phe Glu Ala Gly Ala Ala Ala Gly Ala Ala Pro 92

GGG GAA GAA GAC CTG TGT GCA GCA TTT AAC GTC ATA TGT GAT AAT GTG GGG AAA GAT TGG AGA AGG CTG GCT CGT  
 Gly Glu Glu Asp Leu Cys Ala Ala Phe Asn Val Ile Cys Asp Asn Val Gly Lys Asp Trp Arg Arg Leu Ala Arg 117

500  
 CAG CTC AAA GTC TCA GAC ACC AAG ATC GAC AGC ATC GAG GAC AGA TAC CCC CGC AAC CTG ACA GAG CGT GTG CGC  
 Gln Leu Lys Val Ser Asp Thr Lys Ile Asp Ser Ile Glu Asp Arg Tyr Pro Arg Asn Leu Thr Glu Arg Val Arg 122

600  
 GAG TCA CTG AGA ATC TGG AAG AAC ACA GAG AAG GAG AAC GCA ACA GTG GCC CAC CTG GTG GGG GCT CTC AGG TCC  
 Glu Ser Leu Arg Ile Trp Lys Asn Thr Glu Lys Glu Asn Ala Thr Val Ala His Leu Val Gly Ala Leu Arg Ser 127

700  
 TGC CAG ATG AAC CTG GTG GCT GAC CTG GTA CAA GAG GTT CAG CAG GCC CGT GAC CTC CAG AAC AGG AGT GGG GCC  
 Cys Gln Met Asn Leu Val Ala Asp Leu Val Gln Glu Val Gln Gln Ala Arg Asp Leu Gln Asn Arg Ser Gly Ala

ATG TCC CCG ATG TCA TGG AAC TCA GAC GCA TCT ACC TCC GAA GCG TCC TGATGGGCCGCTGCTTTGCGCTGGTGGACCACAGGC  
 Met Ser Pro Met Ser Trp Asn Ser Asp Ala Ser Thr Ser Glu Ala Ser \*

800  
 ATCTACACAGCCTGGACTTTGGTTCTCTCCAGGAAGGTAGCCCAGCACTGTGAAGACCCAGCAGGAAGCCAGGCTGAGTGAGCCACAGACCACCTGCTTC

900  
 TGAACCTAAGCTGCGTTTATTAATGCCTCTCCCGCACCAGGCCGGGCTTGGGCCCTGCACAGATATTTCCATTTCTTCTCACTATGACACTGAGCAAGA

1000  
 TCTTGTCTCCACTAAATGAGCTCCTGCGGGAGTAGTTGGAAGTTGGAACCGTGTCAGCACAGAAGGAATCTGTGCAGATGAGCAGTCACACTGTTACT

1100  
 CCACAGCGGAGGAGACCAGCTCAGAGGCCAGGAATCGGAGCGAAGCAGAGAGGTGGAGAACTGGGATTGTAACCCCCGCCATCCTTCACCAGAGCCCAT

1200  
 GCTCAACCACTGTGGCGTTCTGCTGCCCCCTGCAGTTGGCAGAAAGGATGTTTTGTCCCATTTCCTTGGAGGCCACCGGGACAGACCTGGACACTAGGGTC

1300  
 AGGCGGGGTGCTGTGGTGGGGAGAGGCATGGCTGGGGTGGGGTGGGGAGACCTGGTTGGCCGTGGTCCAGCTCTTGGCCCCCTGTGTGAGTTGAGTCTCC

1400  
 TCTCTGAGACTGCTAAGTAGGGGAGTGATGGTTGCCAGGACGAATTGAGATAATATCTGTGAGGTGCTGATGAGTGATTGACACACAGCACTCTCTAAA

1500  
 TCTTCCTGTGAGGATTATGGGTCTGCAATTCTACAGTTTCTTACTGTTTTGTATCAAAATCACTATCTTTCTGATAACAGAATTGCCAAGGCAGCGGG

1600  
 ATCTCGTATCTTTAAAAAGCAGTCTCTTATTCCTAAGGTAATCCTATTAAAA

FIGURE 2A

hFADD  
rFas  
hFas  
hTNFR-1

(111-145)  
(217-251)  
(228-262)  
(341-375)

DD A K K F F V R R L A R Q L K V S D T K I D S I E D R Y P R N L T E R V R E S L  
 DD A K K F F V R R L A R Q L K V S D T K I D S I E H N S P Q D A A E Q K I Q Q L L L  
 QQ V K K F F V R R L A R Q L K V S D T K I D S I E H N S P Q D A A E Q K I Q Q L L L  
 RR A K K F F V R R L A R Q L K V S D T K I D S I E H N S P Q D A A E Q K I Q Q L L L

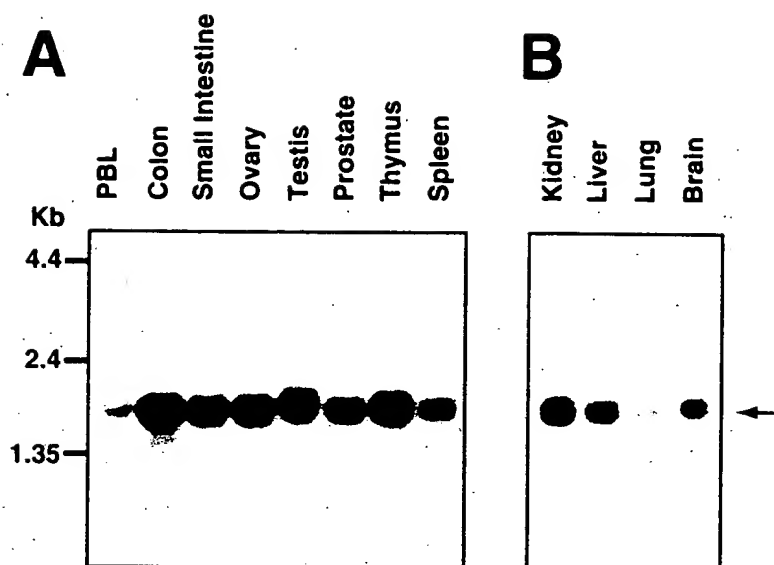


hFADD  
rFas  
hFas  
hTNFR-1

(146-180)  
(252-286)  
(263-297)  
(376-410)

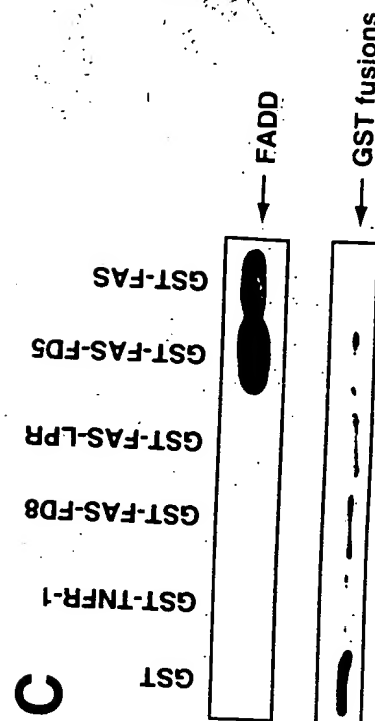
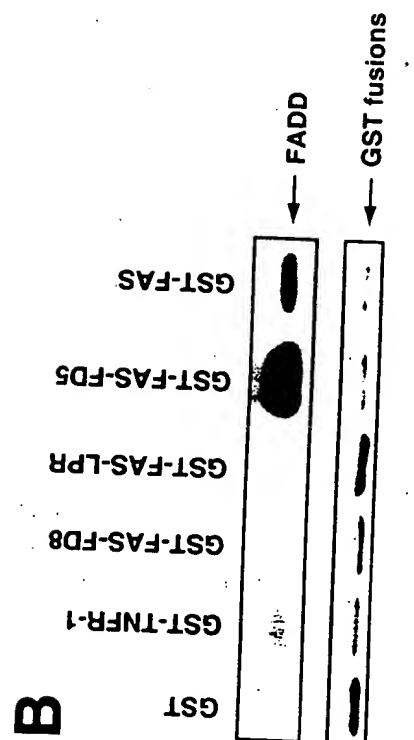
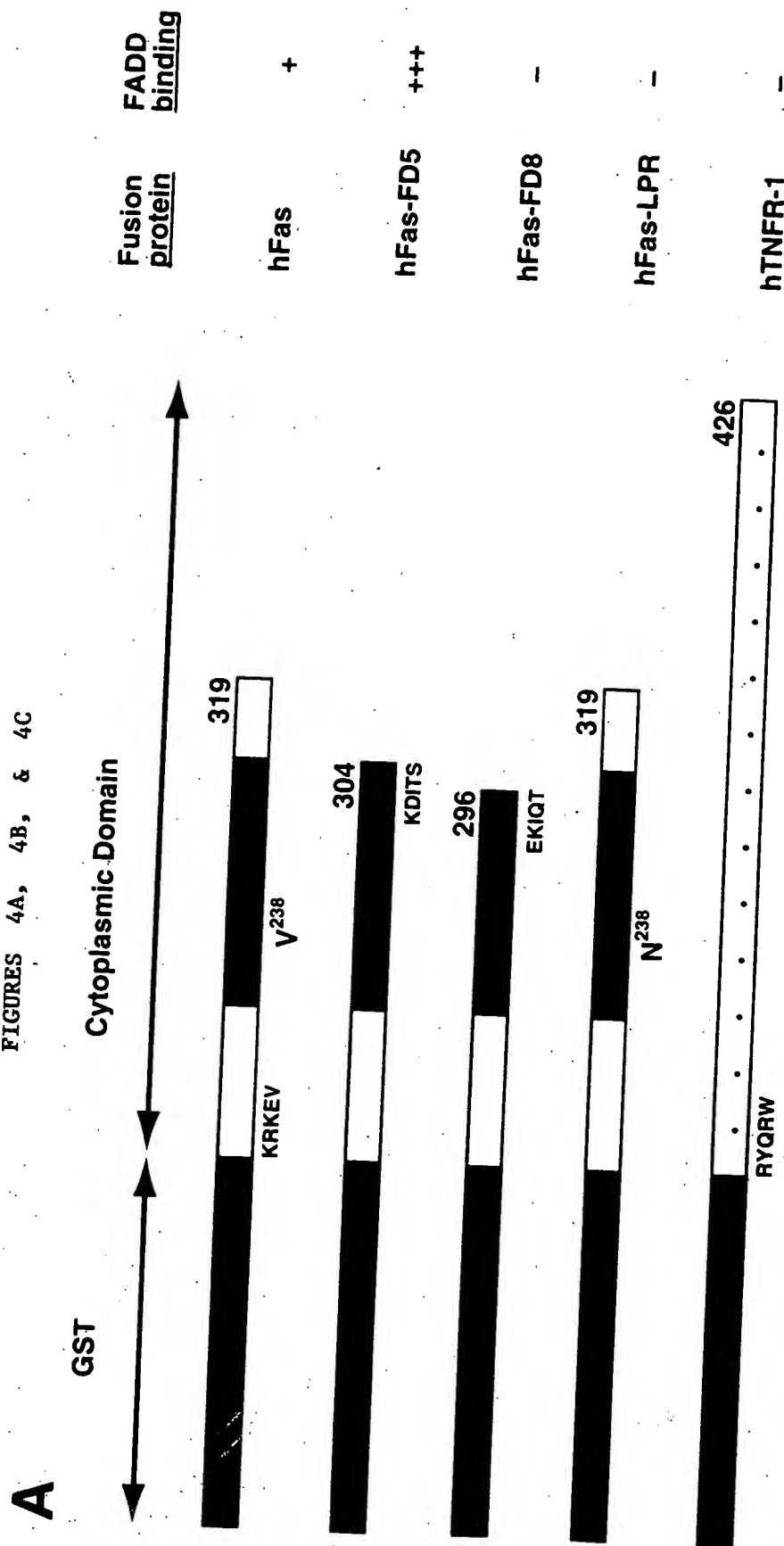
P R T U K E E N A T V A H L V G A L R S C Q M N L V A D L V Q E V  
Q Q C N W R Y Q Q S H G K T G A C Q A L I Q G L R K A N R C D I A E I Q A M  
A W W R R R L P R R K K E A T L E L L C R V L R K K A N R C T L A E I Q T I E

**FIGURE 2B**



FIGURES 3A &amp; 3B

FIGURES 4A, 4B, &amp; 4C



**A**

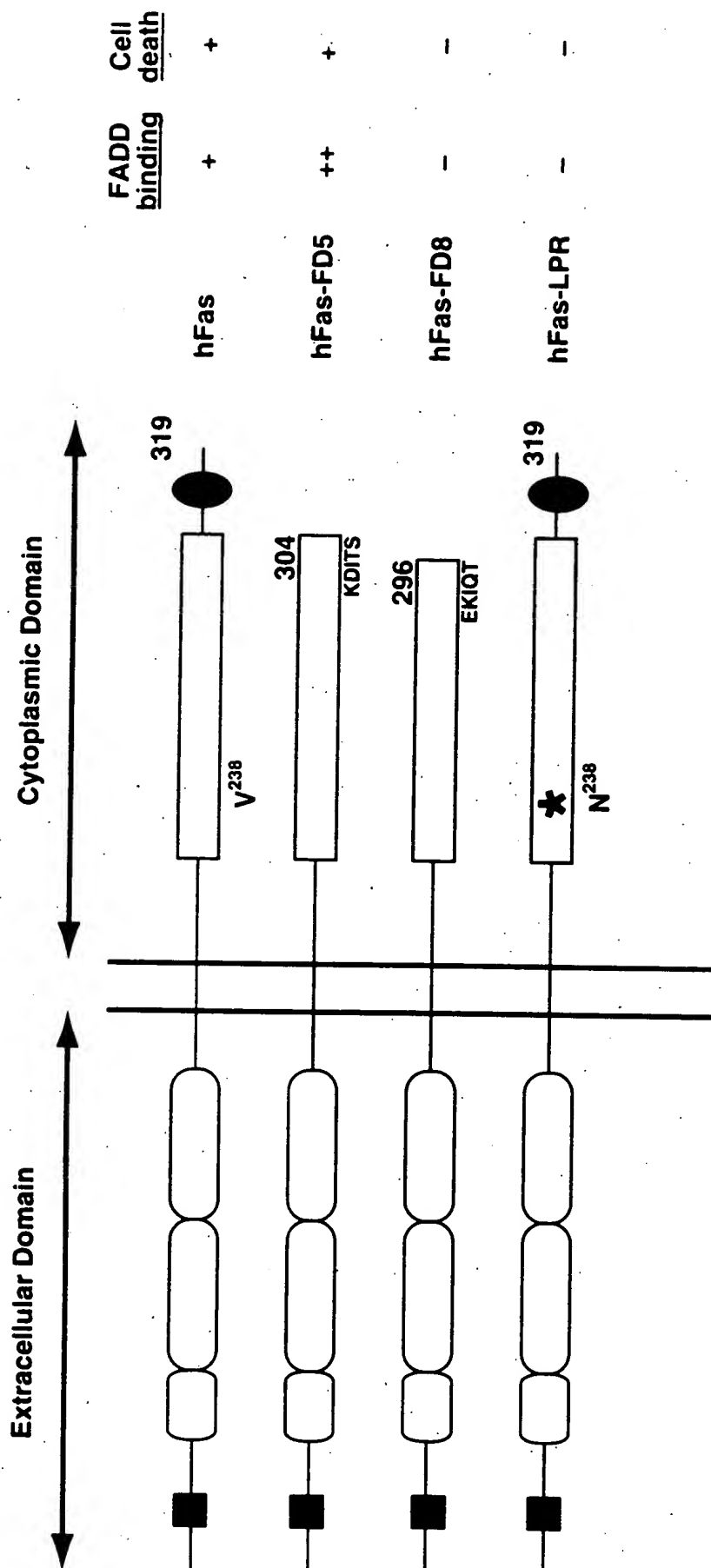
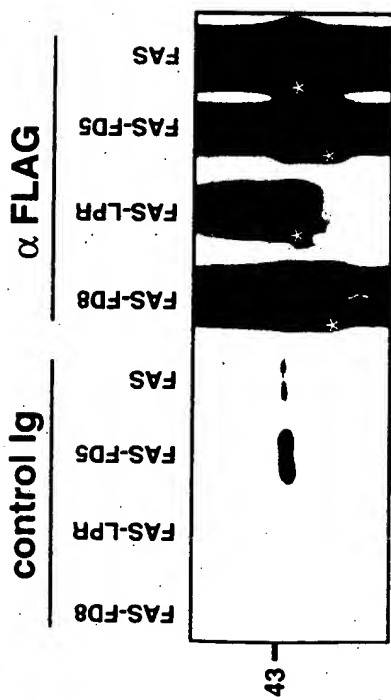
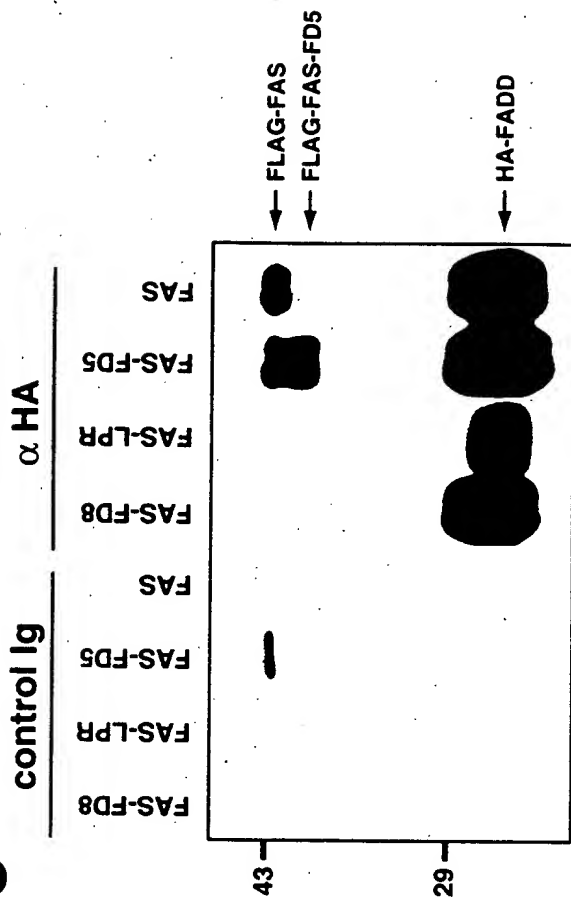
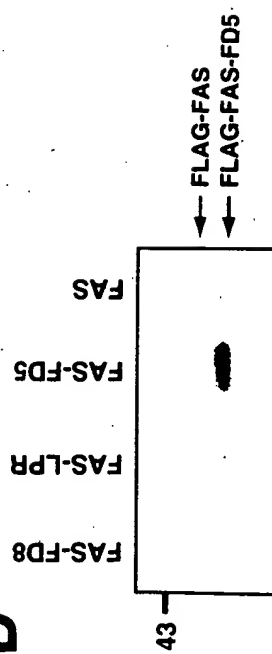


FIGURE 5A

**B****C****D**

FIGURES 5B, 5C, &amp; 5D

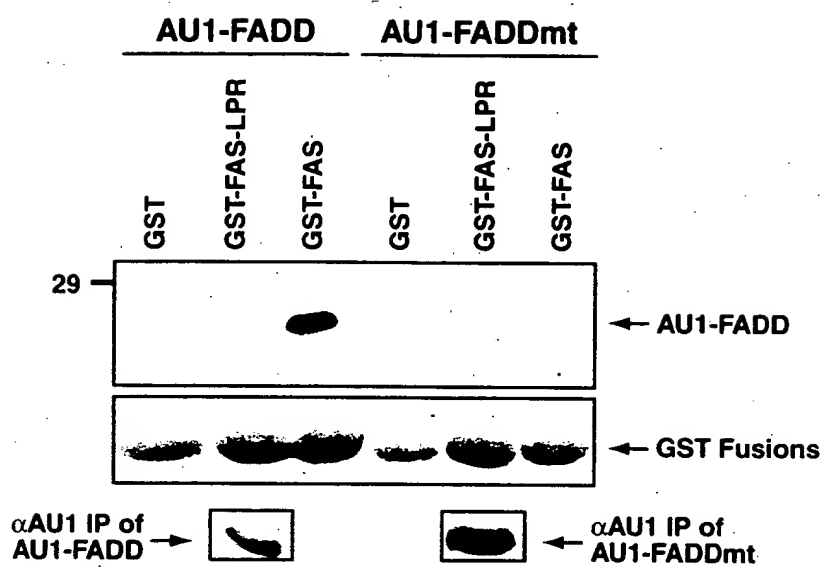
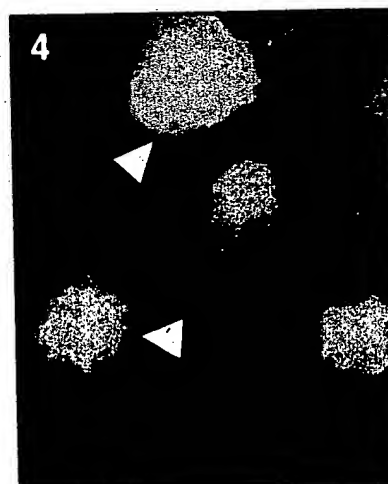
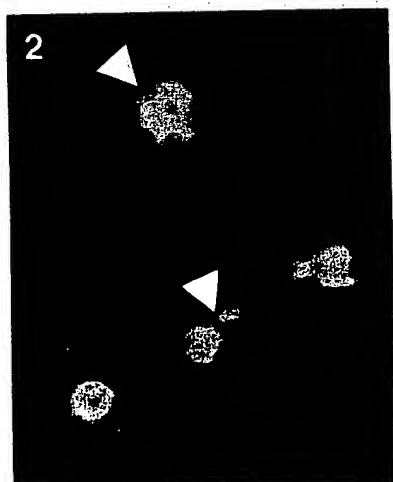
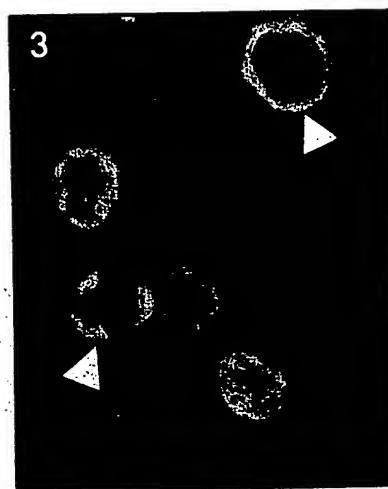
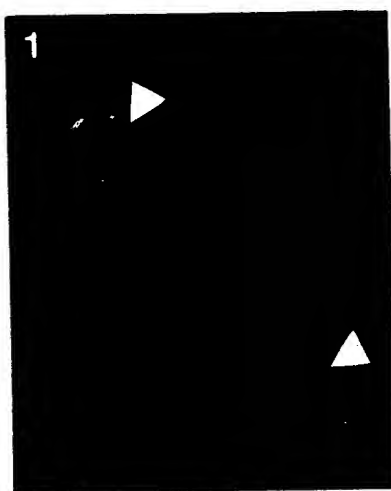


FIGURE 6

**A****FIGURE 7A**

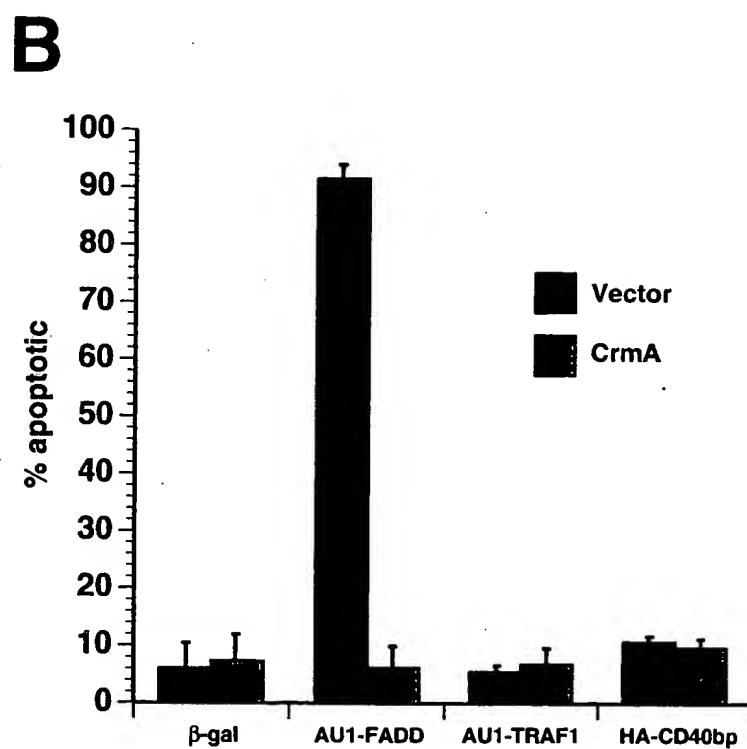


FIGURE 7B

C

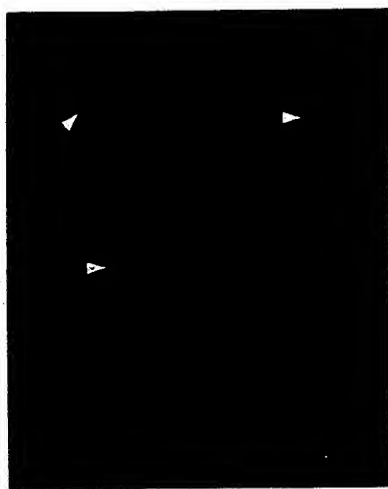


FIGURE 7C

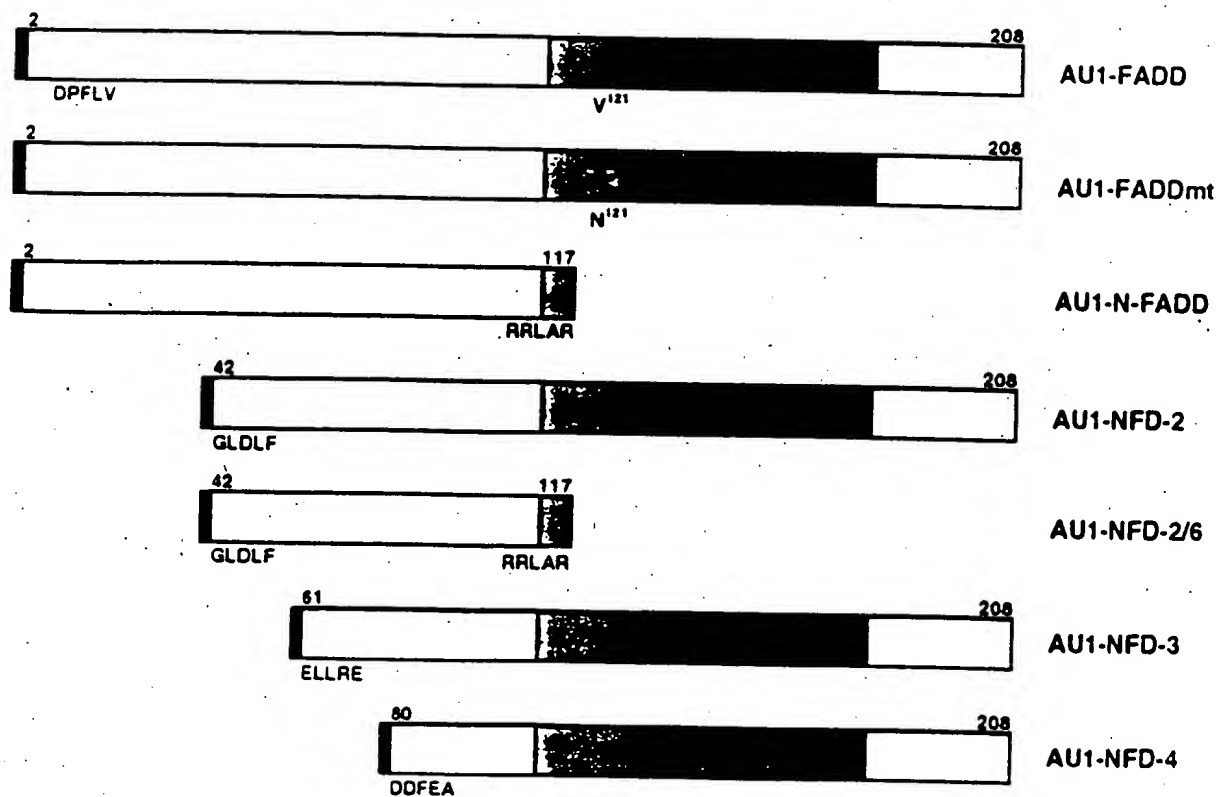
**A****B****Cell  
Death**

FIGURE 8A &amp; 8B